

FIG. 1

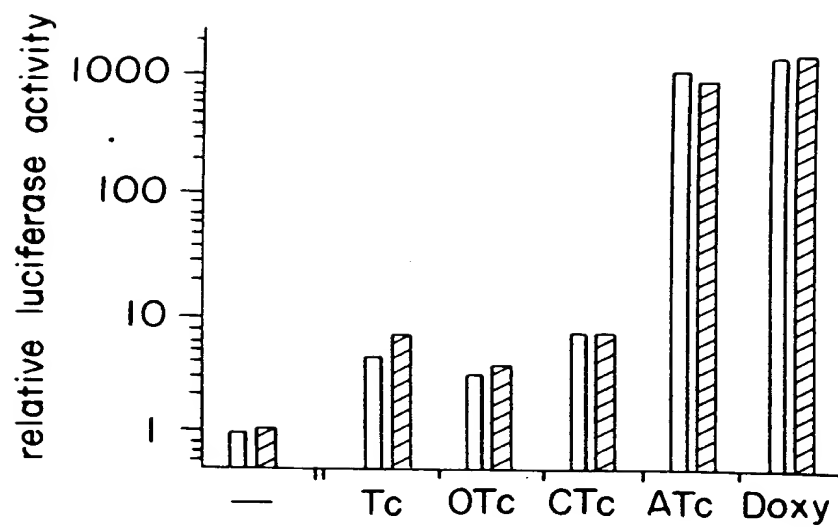


FIG.2

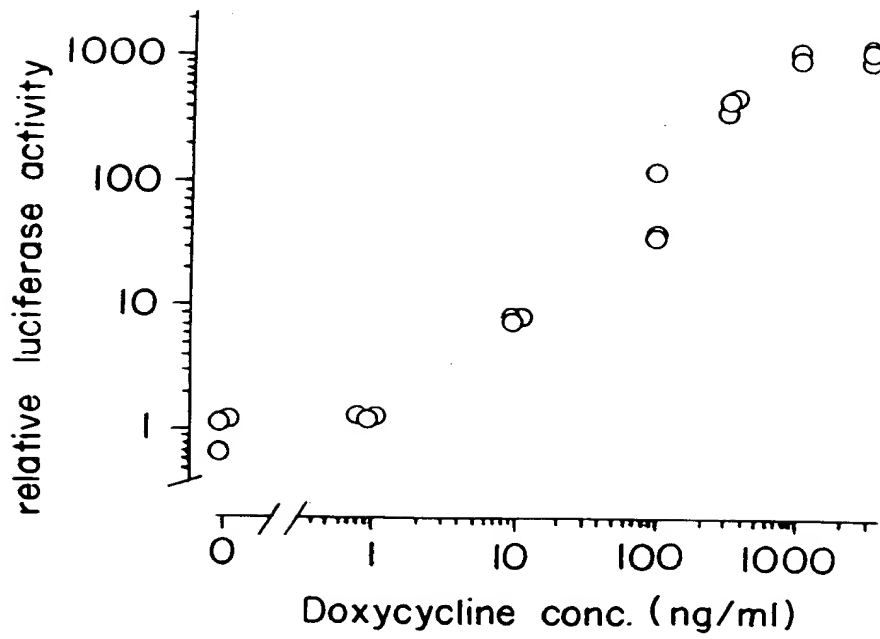


FIG. 3

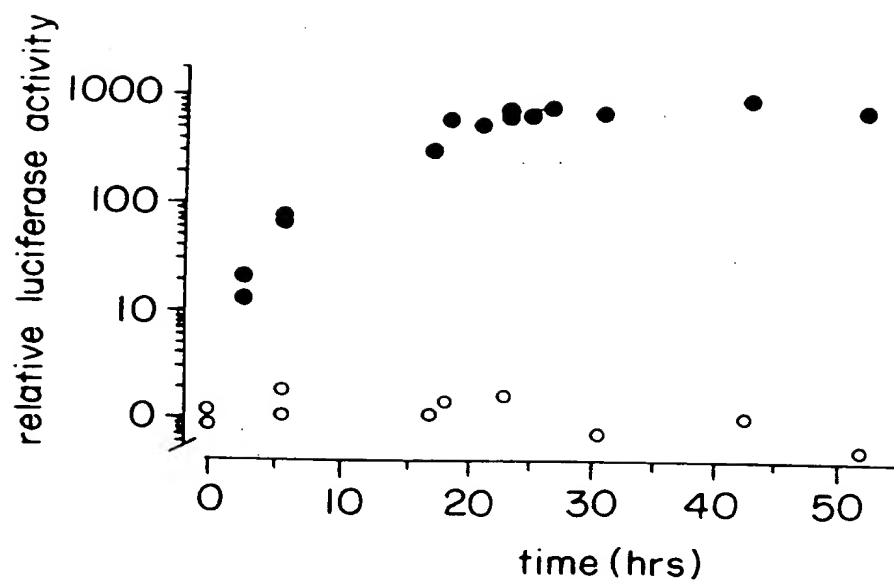
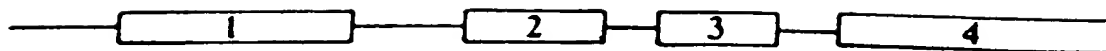


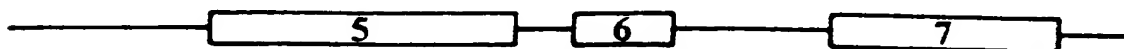
FIG. 4

H T H



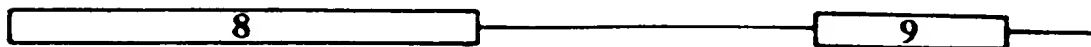
MSRLQSKVINSALELLNEVGIEGLTTRKLAQKLGVEOPTLYWHYKNKRALLDALAIEMLD RH
 MA--NRES--DA--G---T--DE-----I-----V-I-A--
 MTK-QPNT--RA--D-----VD-----ER--Q--A---FR-----EA--AEN
 MNK-QREA--RT--G---D--M-----R--ER--Q--A---F-----EA--TIN
 MTK---GT--AAG-----MDS-----ER-K-Q--A---FQ-----PEA--RER
 MA--SLDD--SM--T--DSE-L-----S-KI-----R--QT-MNM-SEAI-AK-

63 B
D
A
C
G
E



HTHFCPLEGESWQDFLRNNAKSFRCALLSHRDGAKVHLGTRPTEKQYETLENQLAFLCQOGFS
 -DYSL-AA-----S-----M--R--RY-----D---D-V-T--R-MTEN--
 ---SV-RADD--RS--IG--R--O--AY---RI-A---GAP-M--ADA--R--EA--
 ---ST-R0000-RS--KG--C--R--AY---RI-A---AAP-M-KADA--R--DA--
 ---RSL-E-N-D-RV--KE--L--T---Y---RI-A---PNFG-A-T-IR--AE--C
 ---RSA--PT---Q--QE--L--K--V---RL-I--S--PP-F-QA-A--RC--DA--

126 B
D
A
C
G
E



LENALYALSAVGHFTLGCVLEDOEHQVAKEERE TPTTOSMPPLL RQAIELFDHQGA
 -RDG---I---S-----A---Q---TA-LTD-P AAPDENL-----E-LQIM-SDDG
 AGD-VN--MTISY--V-A---E-AGDSESG--GG -VEQAPLS---A--DA--EA-P
 AGD-T---M-ISY--V-A---Q-ASEADA--GEDQL-TSAST--AR-QS-MKIVYEA-P
 PKR-VW--R--S-YVV-S---Q-ASDAD -VPORPDVSEQAPSSF-HVLFHELETD-M
 V-E--FI-QSIS-----A---E-ATNQIENNHV I-AA---QE-FNIQARTS-

182 B
182 D
183 A
186 C
184 G
179 E



EPAFLFGLLELIICGLEKOLKCESGS
 -Q---H---SL-R-F-V--TALLQIVGGOKLIIPFC
 DA--EQ--AV-VD--A--RRLVVRNVEGPRKGDD
 DA--ER--A---G---MRLTTNDIEVLKNVDE
 DA--N---DSL-A-F-RLRAAVLATD
 -M--H---KSL-F-FSA--DEKKHTPIEDGNK

207 B
218 D
216 A
219 C
210 G
211 E

FIG. 5

A1	AC TTTATCA C TGATAAACA TGAAATAGTG A CTAATTGT	AAC TTTATCAG TGATTAAGA TTGAATAGT CACTAATTCT	A2
B1	ACTCTATCA T TGATAGAGT TGAGATAGT A ACTATCTCA	TCCCTATCAG TGATAGAGA AGGATAGT CACTATCTCT	B2
C1	AGCTTATCA TC GATAAGCT TCGAATAGT AGCTATTCCGA	AGTTTAT CACAGT TAAAT T TCAAAAT AGTGTC AATTTA A	C2
D1	ACTCTATCA T TGATAGGGA TGAGATAGT A ACTATCCC T	ACTCTATCA A TGATAGGGA TGAGATAGT T ACTATCCC T	D2
E1	AATCTATCA C TGATAGAGT TTAGATAGT G ACTATCTCA	ACCCTATCA TCGATAGAGA TGGGATAGT AGCTATCTCT	E2

FIG. 6

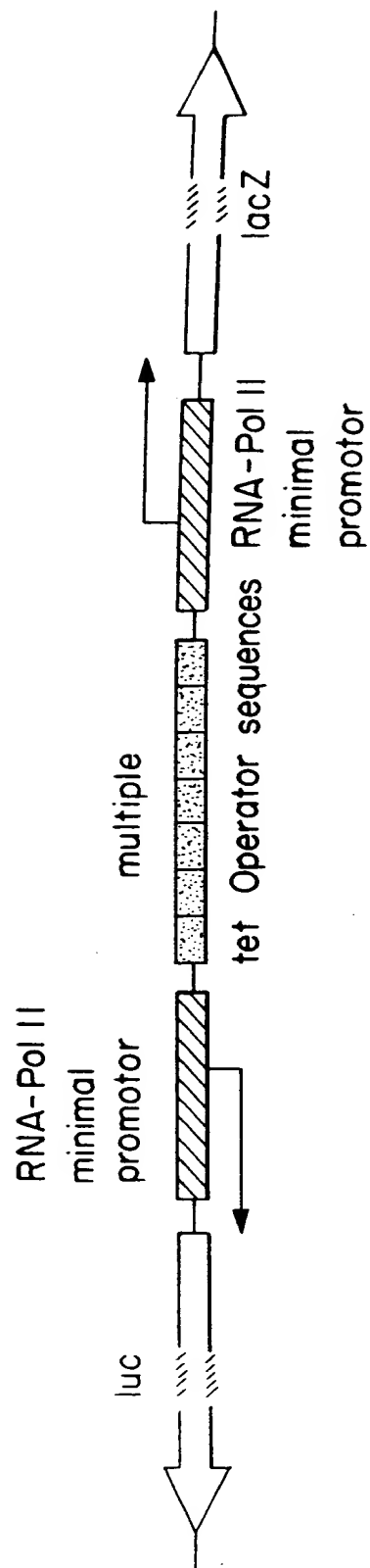


FIG. 7A

5' GAATTCGGGG
EcoRI + 75

CCGCGGAGGCTGGATCGGTCCCGGTGTCTTCTATGGAGGTCAAAACAGCGTGGA

+ 1
←
C

P_{hCMV}*-3

TGGCGTCTCCAGGCGATCTGACGGTTCATAAACGAGCTCTGCTTATATAGG
- 31

tet O

TC (GAGTTACCACTCCCTATCAGTGATAGAGAAAAGTGAAAGTC)₇GAGC

P_{hCMV}*-1

TCGGTACCCGGGTCGAGTAGGCGTGTACGGTGGGAGGCCTATATAAGCAGAG
- 53

CTCGTTTAGTGAACCGTCAGATCGCCTGGAGACGCCATCCACGCTGTTTTGA
+ 1 →

CCTCCATAGAAGACACCGGGACCGATCCAGCCTCCGCGGCCCGAATTC 3'
+ 75 EcoRI

1.34030: 6364466

FIG. 7B

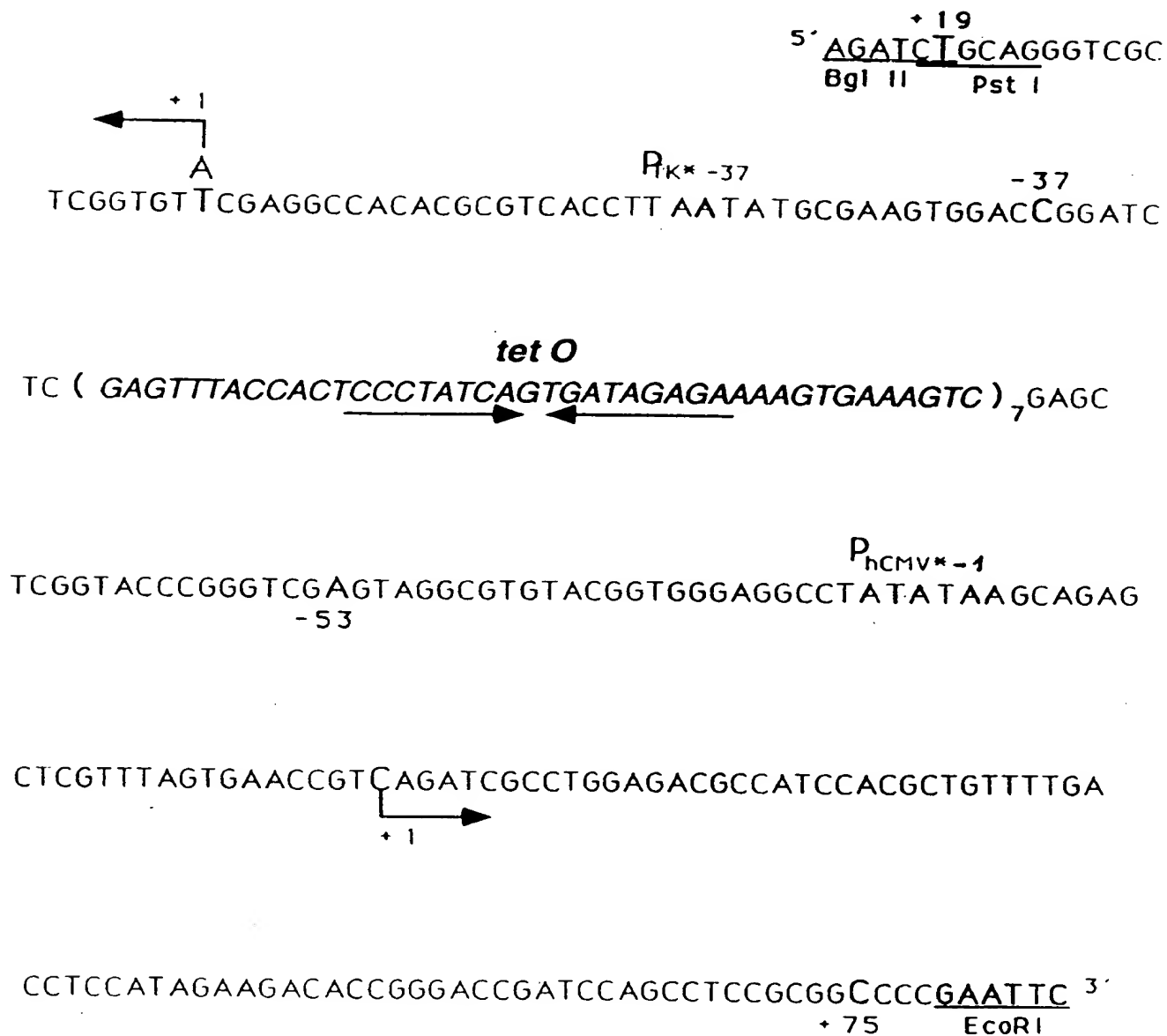


FIG. 8A

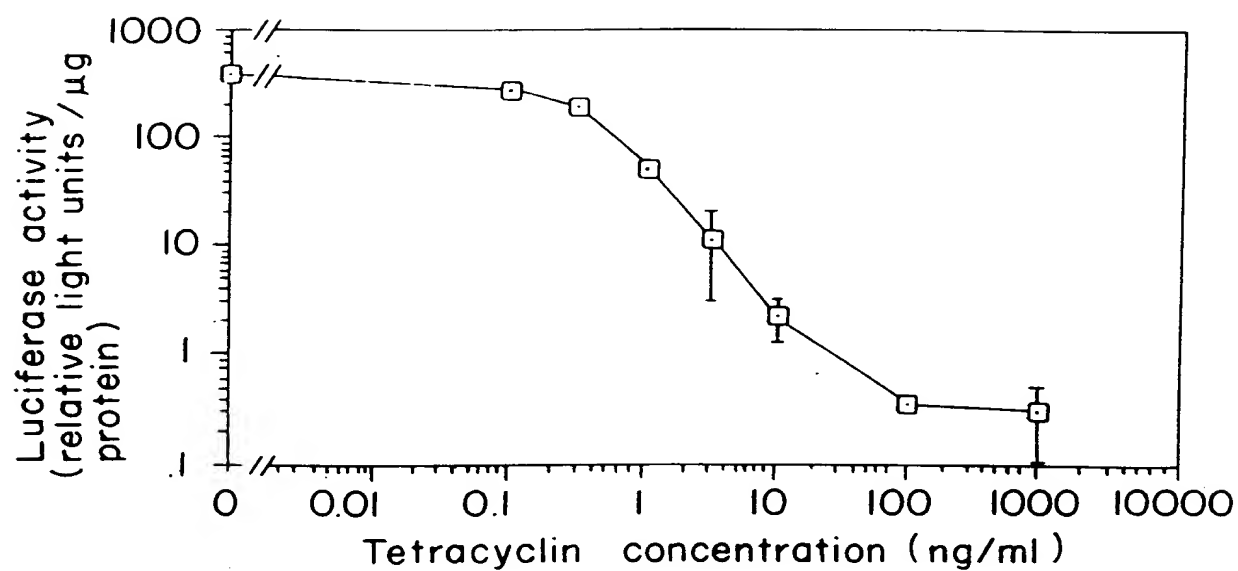


FIG. 8B

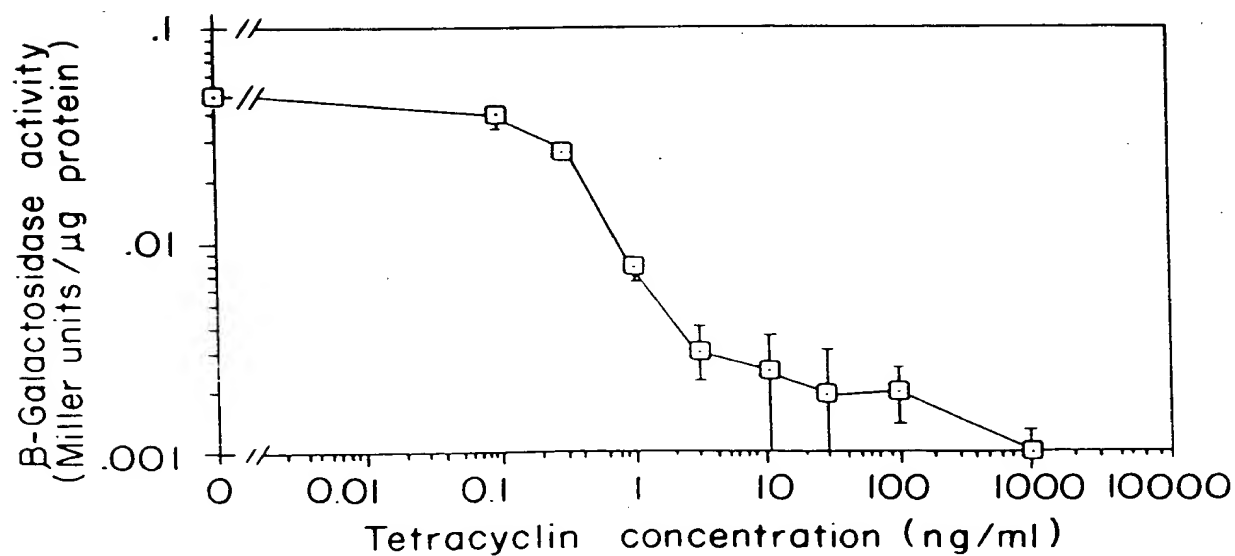


FIG. 9A

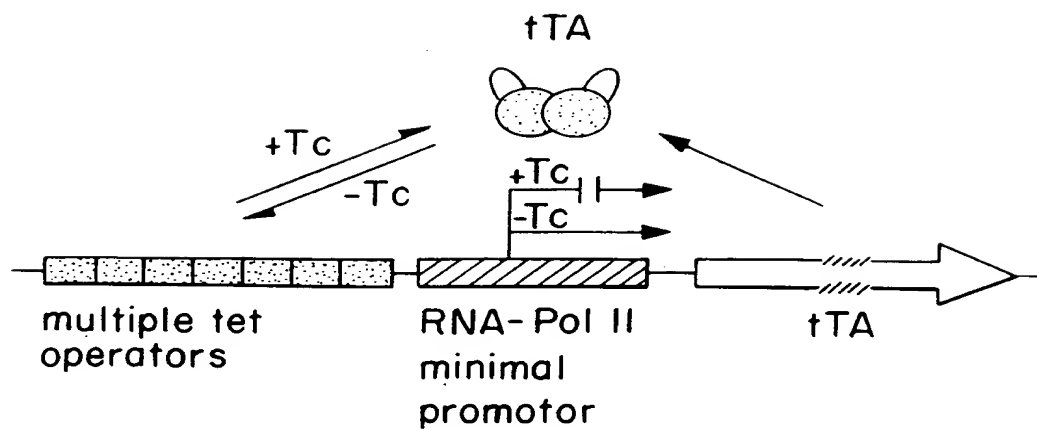


FIG. 9B

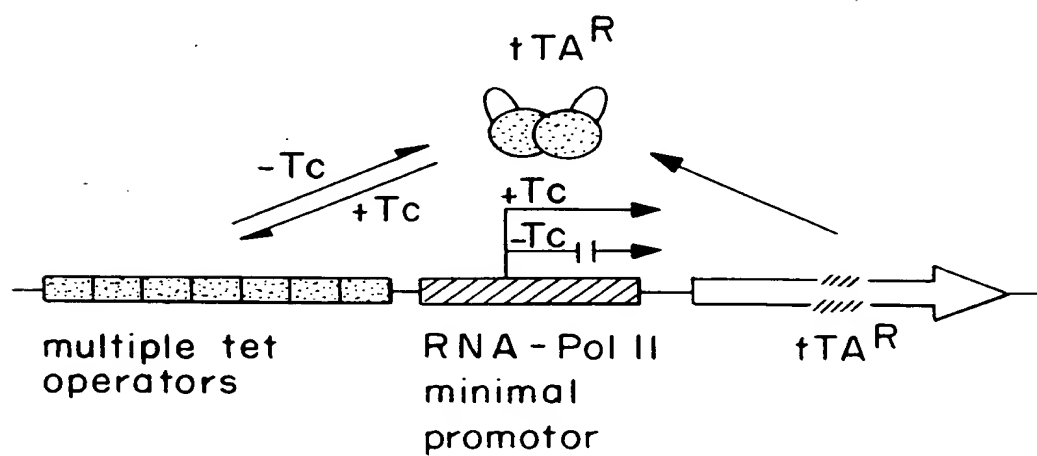


FIG. 10

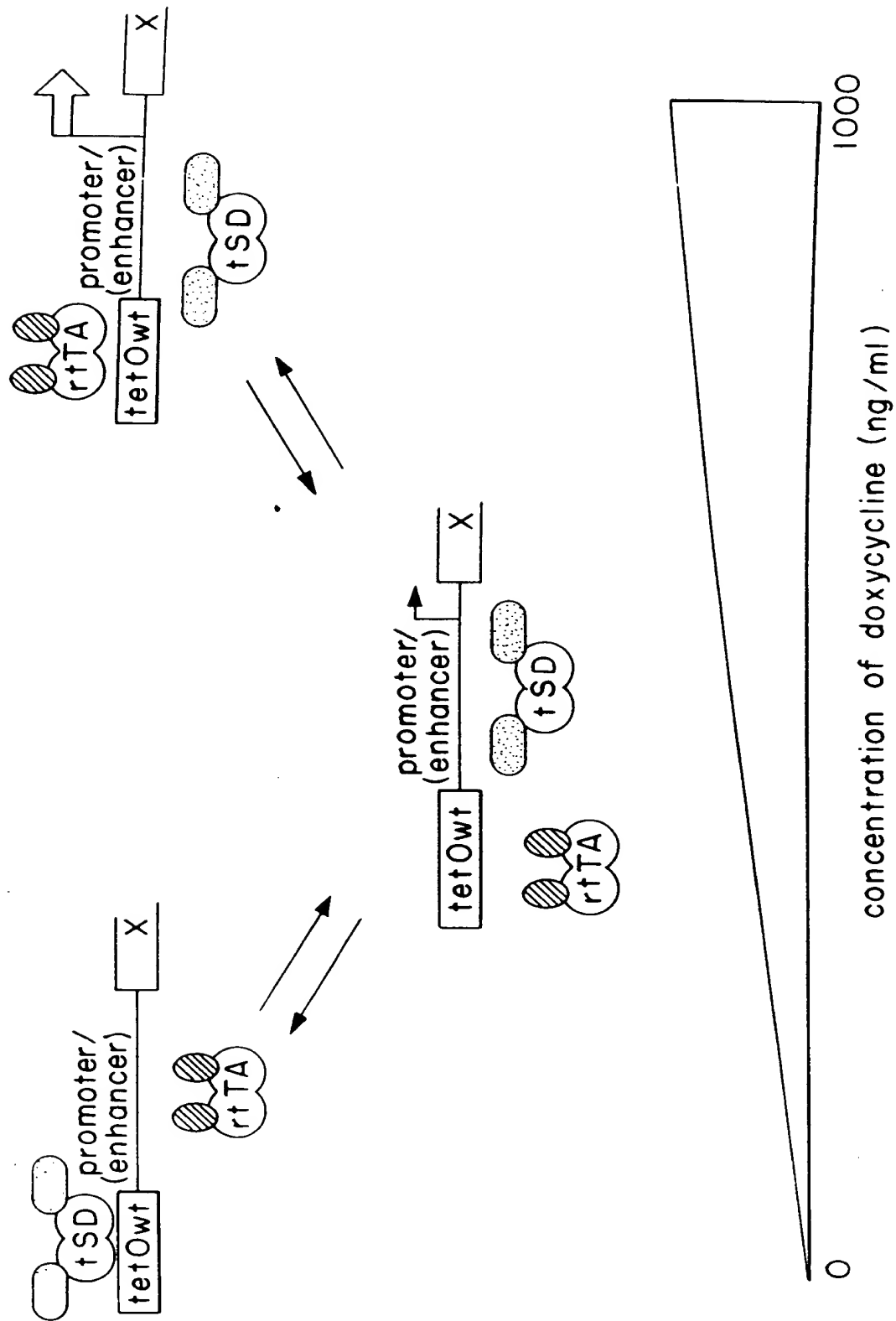


FIG. 11

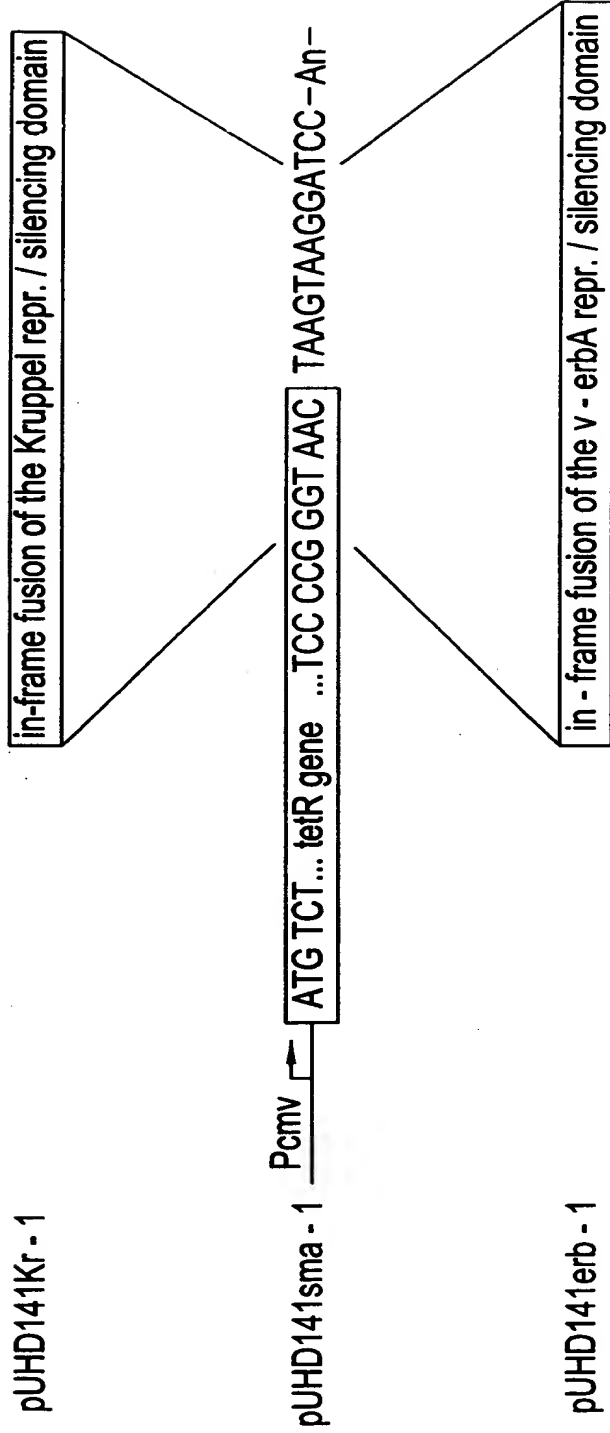


FIG.12

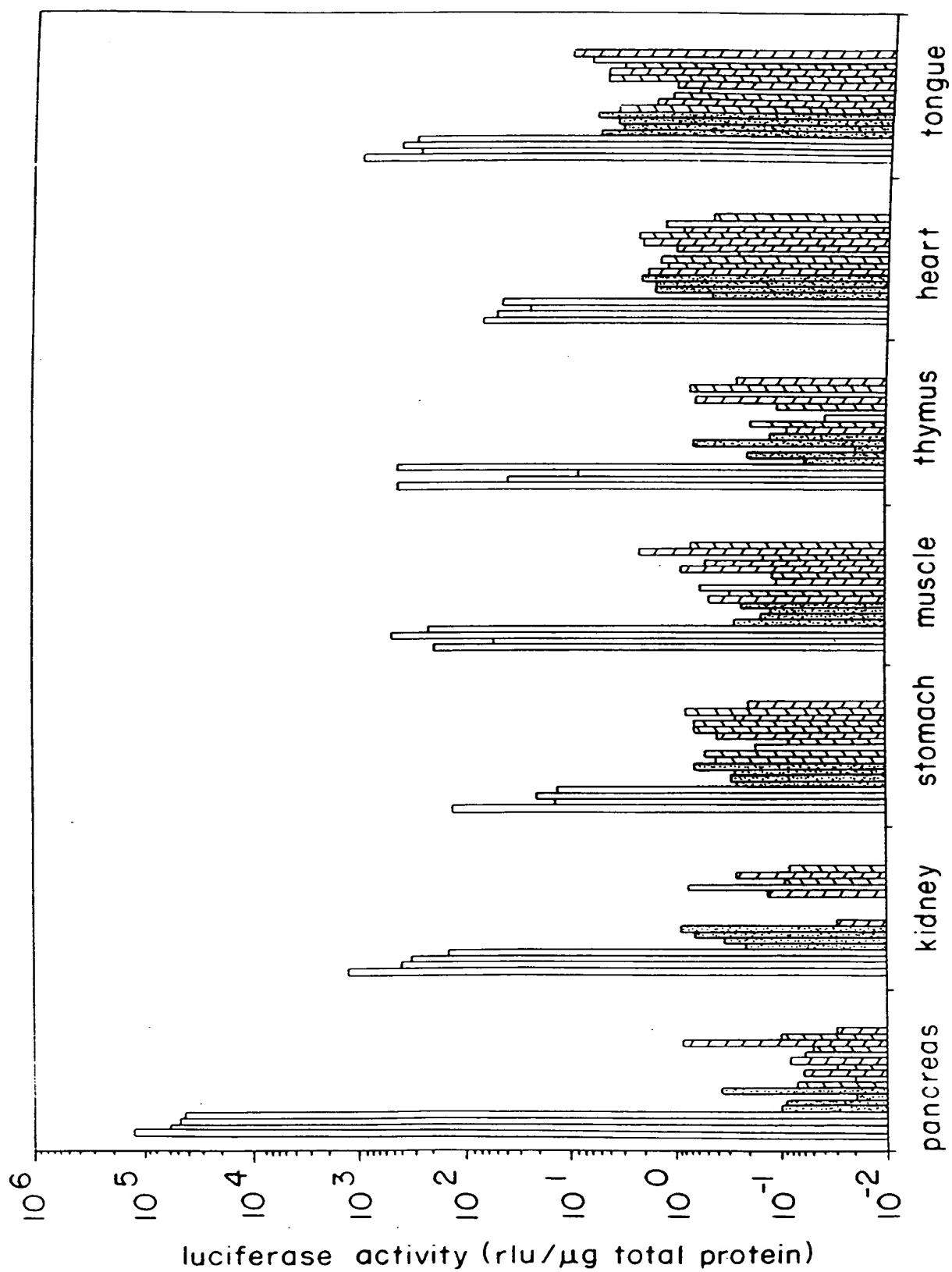


FIG. 13

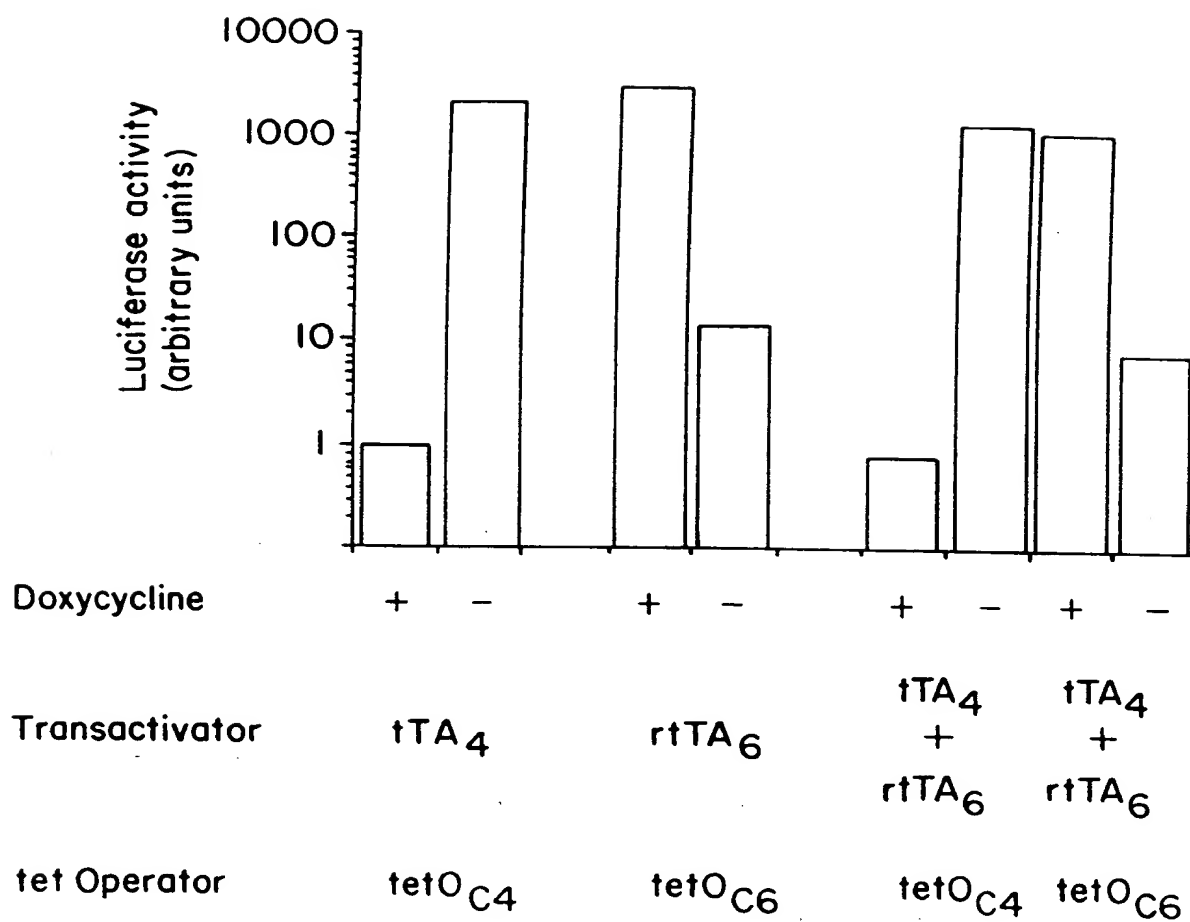


FIG.14A

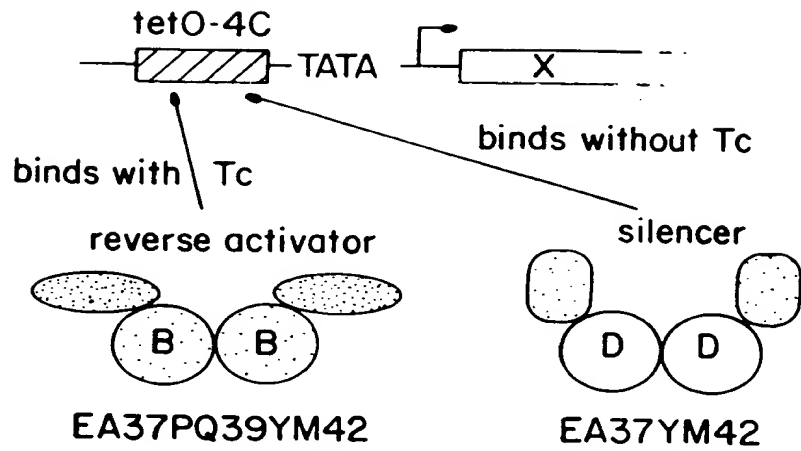


FIG.14B

